| 0 | 1, O | 96 | 144 | 192 | 240 | 288 | 336 | 384 | |
|----------------|------------------------|------------------------|-------------------|------------------|------------------------|------------------------|-------------------|-------------------|-------------|
| E | Val | GAG Glu | TGC Cys | TIC | CGG Arg 80 | AGA Arg | GIC Val | ATT | |
| E | CIT | GGA Gly | ATT Ile | ACT | AAA Lys | GAA Glu 95 | GAA Glu | GGT Gly | |
| { ! | Al'I Ile | AGT Ser | TTT Phe | ACT | ATG Met | CAG Gln | GCT Ala 110 | AAA Lys | |
| | CCA Pro | GTC Val | AAA Lys. 45 | GIC | CAT His | GTA Val | CGT | TTA Leu 125 |) 1 1 |
| | GIC Val | Ser 30 | CII Leu | CTT Let 60 | GAT | $_{\rm TYr}^{\rm TAT}$ | ACA Thr | GAG Glu | |
| | GTT Val | TTT Phe | ACC Thr | ACA Thr | CCA Pro 75 | GGT | AAG Lys | ATC | 14 |
| | GIY Oly | AAA Lys | CTT | CCA | $_{\rm TYr}^{\rm TAC}$ | GAA Glu | TAC | AGA Arg | FIG. |
| | ACT | CAC His | AAA Lys | IGG | AGA Arg | CCC | AAC Asn 105 | AAT Asn | |
| | TIC | $_{\rm GIY}^{\rm GGG}$ | GLY | CCA Pro | TCA | ATG Met | 666 61y | GTT Val | T Z O |
| | CIT | AAT Asn | \mathtt{TAC} | GTT Val | TTT Phe | GCC | GAC Asp | CIT | |
| | GAA Glu | GTT | ACA Thr | CCT | TGC | AGT Ser | GAT | ACC | |
| | GAA Glu | GAT Asp | GCA Ala | CTA | CAA | AAG Lys | 85 AAA Lys | GAT | |
| | $_{\rm GLY}^{\rm GGA}$ | GGT | GAT ASP | AAA Lys | GTT Val | TTC | TTC Phe | | ı |
| NO:1: NO:2: | AAA Lys | GAT Asp | GGT | 35 GGA GLY | GGT Gly | TTT Phe | TTT Phe | GAA Glu | |
| | AGT Ser | TTA | GAA Glu | ACT | 5U TAT TYT | · • | | TTT Phe | |
| SEO | ATG Met | I GAA Glu | GGT | ACT | TCT | 65 CAT His | ACT | AAG Lvs | 7 |
| | | | | | | | | | |

| HOOMYWOM CARROL | | |
|-----------------|----------|-------|
| YUDU DAU | | 1 |
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| 432 | 480 | 228 | 576 | 526 | 672 | 716 |
|----------------------------------|--|-------------------|-------------------|---------------------|-------------------|-------------------|
| AAC Asn | 6678 617 160 | GTT Val | CCT | Ser | GTA Val | |
| TAC | AAT Asn | AGC Ser 175 | GGC Gly | CIT | TTT | TA |
| GAA Glu | AAG Lys | GGA Gly | GAT Asp 190 | GCC Ala | GAG Glu | AAA Lys |
| HAG Lea | CAA Gln | GAT | 990 617 | ACT SOFT 2057 | CIT | TAC Tyr |
| AAA Lys 140 | AAA Lys | GAA Glu | ATT 110 | CAA Gln | CIT Leu 220 | CIA |
| CAC | GAC ASP 155 | ATT Ile | OCA Pro | ACA Thr | GIC Val | GAA Glu 235 |
| GGA Gly | GCA Ala | AAC Asn 170 | ACT | Ser | ATG Met | GAT Asp |
| CIT | ATG Met | CAC His | AAT Asn 185 | CIG | CAC His | ATG Met |
| ATT Ile | ATC 11e | AGA Arg | GAA | TAC TVC 200 | GAC | 990 91√ 91√ |
| AAC Asn 135 | TAC | ATT | CAA | CAT His | AGA 215 | CAT His |
| $\frac{\text{GGA}}{\text{Gl} Y}$ | GTA Val 150 | | TAT Tyr | AAC Asn | AAG Lys | ACA Thr 230 |
| GAT | AAT Asn | TTC Phe 165 | CAT His | GAC | GAA Glu | ATT |
| GAA Glu | CAC His | AAC Asn | GAC 180 180 | | AAC | 666 617 |
| AAA Lys | TCA | GIT | GCA Ala | TTA Leu 195 | | GCT |
| TTT Phe 130 | | AAA Lys | CIA | CIT | GAT ASD 210 | |
| GAT | 11 A T 14 A T 14 A T 15 | ATC | CAA | GTC Val | AAA Lys | ACA Thr 225 |

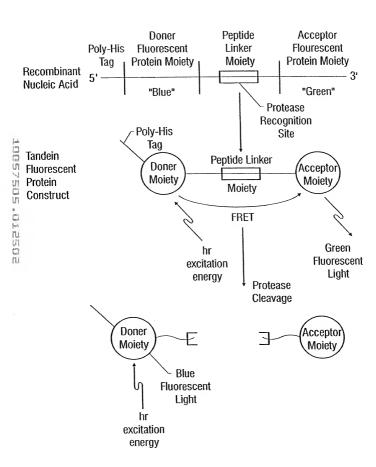
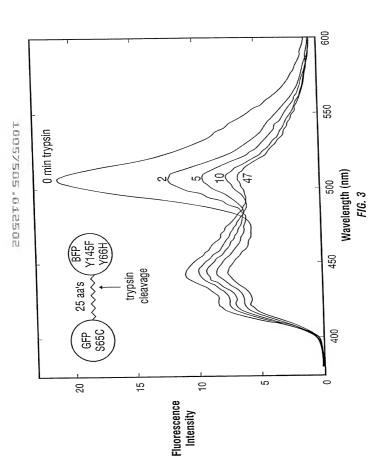


FIG. 2



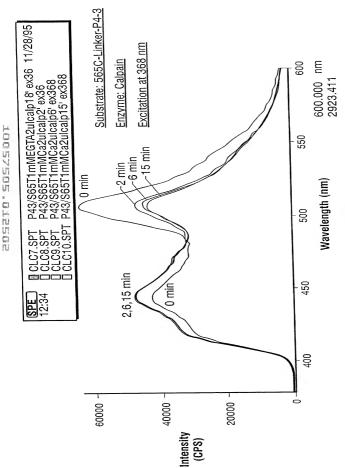
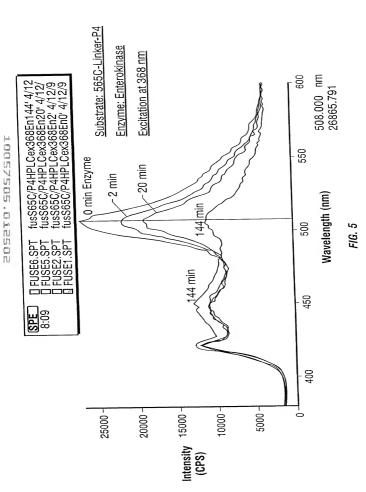
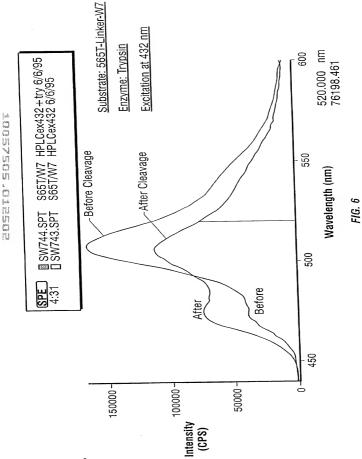
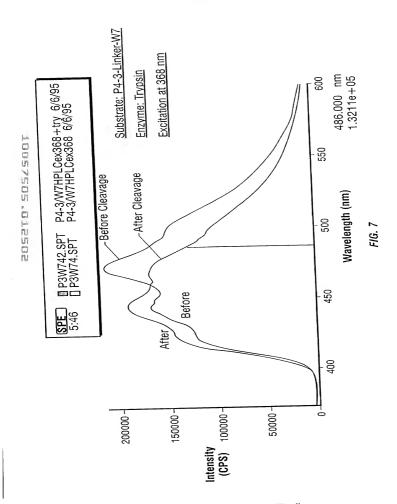


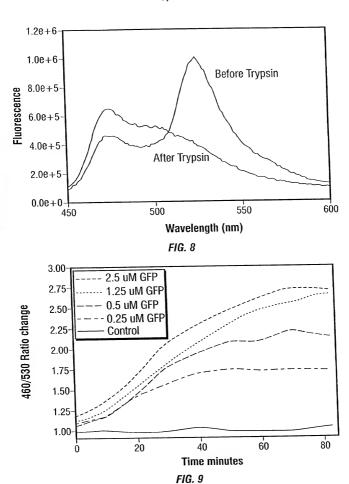
FIG. 4



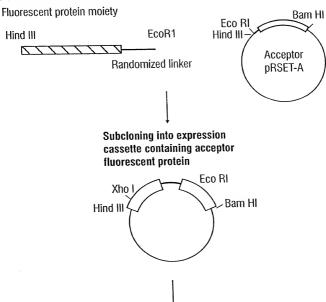








PCR to add randomized linker



Transfection and Screening in cells to identify sequences that are cleaved when co-expressed with the target enzyme